SWLA1: LIGHT CHAIN SEQUENCE DNA AND AMINO ACID SEQUENCE OF THE VL DOMAIN OF CHIMERIC ANTIBODY TEDW EcoRV (242) GGGGATATCCACCATGGAGACAGACACTCCTGCTATGGGTGCTGCTGTTCTGGGTTCCAGGTTCCACAGGTGACATTGT ▶ M E T D T L L L W V L L L W V P G S T G D I V Pstl (377) GCTGACCCAATCTCCAGTTTCTTTGGCTGTCTCTAGGGCAGAGGCCACCATATCCTGCAGAGCCAGTGAAAGTGTTGA L T Q S P V S L A V S L G Q R A T I S C R A S E S V D Kpnl (427) TAGTTATGGCAATAGTTTTATGAACTGGTACCAGCAGAAACCAGGACAGCCACCCCCAACTCCTCATCTATCGTGCATCCAA SYGNSFMNWYQQKPGQPPQLLIYRASN Xbal (482) TCTAGAATACGGGATCCCTGCCAGGTTCAGTGGCAGTGGGTCTAGGACAGACTTCACCCTCACCATTAATCCTGTGGAGGC LEYGIPARFSGSGSRTDFTLT!NPVEA TGATGATGTTGCAACCTATTACTGTCAGCAAAATAATGCGGATCCTCCCACGTTCGGAGGGGGGACCAAGTTGGAAATCAA DDVATYYCQQNNADPPTFGGGTKLEIK Sall (650)

B SWLA1: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VH DOMAIN OF CHIMERIC ANTIBODY TEDW

EcoRV (242)

ACGTAAGTCGACGCT

RKS

GGGGATATCCACCATGGCTGTCTTGGGGCTGCTCTTCTGCCTGGTGACATTCCCAAGCTGTGTCCTGTCCCAGGTGC

M A V L G L L F C L V T F P S C V L S Q V

AGCTGAAGGAGTCAGGACCTGGCCTGGTGGCGCCCTCACAGAGCCTGTCCATCACATGCACTGTCTCAGGGTTCTCA

Q L K E S G P G L V A P S Q S L S I T C T V S G F S

TTAACCAACTATGATATAAATTGGGTTCGCCAGCCTCCAGGAAAGGGTCTGGAGTGGCTGGGAATAATATGGGTGA

L T N Y D I N W V R Q P P G K G L E W L G I I W G D

CGGGAGCACAAATTATCATTCAGCTCTCATATCCAGACTGAGCATCAGCAAGGATAACTCCCAAGAGCCAAATTTTCT

G S T N Y H S A L I S R L S I S K D N S K S Q I F

TAAAACTGAACAGTCTGCAAACTGATGACACAGCCACGTACTACTGTAACTACCCGTGTTTATATTTCTATGGTATG

L K L N S L Q T D D T A T Y Y C N Y P C L Y F Y G M

Nhel (663) Sall (684)

GACTACTGGGGTCAAGGAACCTCAGTCACCGTCTCTTCAGCTAGCACAACAGCCCCATCAGTCGACCCA

D Y W G Q G T S V T V S S A S

A

SWLA2: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VL DOMAIN OF CHIMERIC ANTIBODY TEFE

EcoRV (243)

 $\tt GGGGATATCCACCATGGATTTTCAGGTGCAGATTTTCAGCTTCCTGCTAATCAGTGTCACAGTCATATTGACCAATGGAGAAA$

▶ M D F Q V Q I F S F L L I S V T V I L T N G E

BstEll (372) Pstl (384)

TTTTGCTCACCCCGTCTCCAGCAATCATAGCTGCATCTCCTGGGGAAAAGGTCACCATCACCTGCAGTGCCAGCTCAAGTGTT

I L L T P S P A I I A A S P G E K V T I T C S A S S S V

Koni (419)

AGTTACATGAACTGGTACCAGCAGAAACCAGGATCTTCCCCCCAAAATCTGGATTTATGGTGTATCCAACCTGGCTTCTGGAGT

S Y M N W Y Q Q K P G S S P K I W I Y G V S N L A S G V

TCCTGCTCGCTTCAGTGGCAGTGGGTCTGGGACATCTTTCTCTTTCACAATCAACAGCATGGAGGCTGAAGATGTTGCCACTT

P A R F S G S G S G T S F S F T I N S M E A E D V A T

ATTACTGTCAGCAAAGGAGTAGTTACCCATTCACGTTCGGCTCGGGGACCAAGCTGGAAATAAAACGTAAGTCGACGCT

R SWLA2: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VH DOMAIN OF CHIMERIC ANTIBODY TEFE

EcoRV (242)

Ndel (295)

AATAAGTACTATAACACAGTCCTGAAGAGCCGGCTCACAATCTCCAAGGATACCTCCAACAACCAGGTATTCCTCAAGAT

N K Y Y N T V L K S R L T I S K D T S N N Q V F L K I

CGCCAGTGTGGACACTGCAGATACTGCCACATACTACTGTGCGCGAATAGAGGGGGGGCTCGGGCTACGATGTTATGGACT

A S V D T A D T A T Y Y C A R I E G G S G Y D V M D

Nhel (675) Sall (696)

ACTGGGGTCAAGGAATCTCAGTCACCGTCTCTTCAGCTAGCACAACACCCCCCATCTGTCGACCCA

Y W G Q G I S V T V S S A S

SWLA3: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE VL DOMAIN OF CHIMERIC ANTIBODY TEFC

EcoRV (242)

GGGATATCCACCATGATGAGTCCTGCCCAGTTCCTGTTTCTGTTAGTGCTCTGGATTCGGGAAACCAACGGTGATGTTGTG MM SPAQFLFLL V L W I R E T N G D V V BstEl (347)

ATGACCCAGACTCCACTCACTTGTCGGTTACCATTGGACAACCAGCCTCCATCTCTTGCAAGTCAAGTCAAGTCAGGCCTCTTA MTQTPLTLSVTIGQPASISCKSSQSLL ▶ D R D G R T Y L S W L L Q R P G Q S P K R L I Y L V S AAACTGGACTCTGGAGTCCCTGACAGGTTCACTGGCAGTGGATCAGGGACAGATTTCACACTGAAAATCAGCAGAGTGGAG K L D S G V P D R F T G S G S G T D F T L K I S R V E GCTGAGGATTTGGGGAGTTTATTATTGCTGGCAAGGTACACATTTTCCGCTCACGTTCGGTGCTGGGACCAAGCTGGAGCTG *A E D L G V Y Y C W Q G T H F P L T F G A G T K L E L Sal (653) AAACGTAAGTCGACC

▶ K R K S

SWLA3: HEAVY CHAIN SEQUENCE B

DNA AND AMINO ACID SEQUENCE OF THE VH DOMAIN OF CHIMERIC ANTIBODY TEFC

EcoRV (1425)

GATATCCACCATGGACITCGGGTTGAGCTTGGTTTTCCTTGTCCTTACTTTAAAAGGTGTCCAGTGTGAAGCTGGT ▶ M D F G L S L V F L V L T L K G V Q C D V K L V GGAGTCTGGGGGAGGCTTAGTGAACCCTGGAGGGTCCCTGAAACTCTCCTGTGCAGCCTCTGGATTCACTTTCAGTAGCTA ESGGGLVNPGGSLKLSCAASGFTFSSY BspEl (1611)

TACCATGTCTTGGGTTGGCCAGACTCCGGAGAGGGCCTGGAGTGGGTCGCATCCATTAGTAGTGGTGGTACTTACACCTA TMSWVRQTPEKRLEWVASISSGGTYTY CTATCCAGACAGTGTGAAGGGCCGATTCACCATCTCCAGAGACAATGCCAAGAACACCCTGTACCTGCAAATGACCAGTCT YPDSVKGRFTISRDNAKNTLYLQMTSL GAAGTCTGAGGACACCCATGTATTACTGTTCAAGAGATGACGCTCCTACGCTCCTATTACTATGCTATCGACTACTG K S E D T A M Y Y C S R D D G S Y G S Y Y Y A M D Y W Nhel (1861)

3

GGGTCAAGGAACCTCAGTCACOGTCTCTTCAGCTAGCTCAACACCCCCATCAGTCGACCCA GQGTSVTVSSAS

SWLA1: LIGHT CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE ABERRANT VL DOMAIN

EcoRV FcoRI. ▶ M E T D T L L L W V L L L W V P G TCCACTGGTGACATTGTGCTGACACAGTCTCCTGCTTCCTTAGCTGTATCTCTGGGGCAGAGGGCCACCATCTCATAC ▶ STGDIVLTQSPASLAVSLGQRATISY AGGGCCAGCAAAAGTGTCAGTACATCTGGCTATAGTTATATGCACTGGAACCAACAGAAACCAGGACAGCCACCCAGA FRASKSVSTSGYSYMHWNQQKPGQPPR EcoO1091 CTCCTCATCTATCTTGTATCCAACCTAGAATCTGGGGTCCCTGCCAGGTTCAGTGGCAGTGGGTCTGGGACAGACTTC ▶ L L I Y L V S N L E S G V P A R F S G S G S G T D F PfiMi ${\tt ACCCTCAACATCCATCCTGTGGAGGAGGAGGATGCTGCAACCTATTACTGTCAGCACATTAGGGAGCTTACACGTTCG}$ T L N I H P V E E E D A A T Y Y C Q H ! R E L T R S GAGGGGGGACCAAGCTGGAAATAAAACGGNCTNATGCTGCACCAACTGTATCCATCTTNAAAANCATCAGTTCTAGAG ▶ E G G P S W K • EcoRI AAGGGCGAATTCC

FIG. 5

SWLA1: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE NON-EFFECTIVE 2ND VH DOMAIN

EcoRV (242)

CIGGICOGCITACIGGGGCCAAGGGACIGIGGICACIGICICTICAGCIAGCACAACACCCCCATCAGICIACCCA

W S A Y W G Q G T V V T V S S A S

SWLA1: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE ABERRANT VH DOMAIN

E∞RV E∞RI ▶ M E T D T L L L W V L L L W V P G STGDIVLTQSPASLAVSLGQRATISY FRASKSVSTSGYSYMHWNQQKPGQPPR EcoO109I $\tt CTCCTCATCTTGTTCCAACCTAGAATCTGGGGTCCCTGCCAGGTTCAGTGGCAGTGGGTCTGGGACAGACTTC$ LLIYLVSNLESGVPARFSGSGSGTDF T L N I H P V E E E D A A T Y Y C Q H I R E L T R S GAGGGGGGACCAAGCTGGAAATAAAACGGNCTNATGCTGCACCAACTGTATCCATCTTNAAAANCATCAGTTCTAGAG ▶ E G G P S W K • **EcoRI** AAGGGCGAATTCC

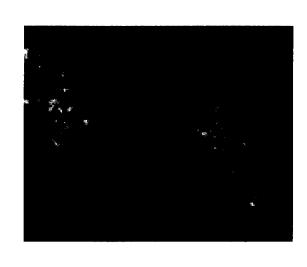
FIG. 7

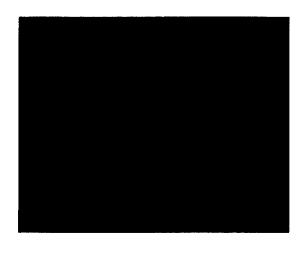
SWLA2: HEAVY CHAIN SEQUENCE

DNA AND AMINO ACID SEQUENCE OF THE ABERRANT VH DOMAIN

EcoRI EcoRV GGAATTCGCCCTTGGGGATATCCACCATGGGATGGACTGGGTCATGCTCTTTCTCCTGGCAGGAACTGCAGGTGTCCT ▶ M G W S W V M L F L L A G T A G V L EcoRV CTCTGAGGTCCAGCTGCAACAGTCTGGACCTGAGCTGGTGAAGCCTGGGGCTTCAGTGAAGATATCCTGCAAGACTTCT SEVQLQQSGPELVKPGASVKISCKTS ${\tt GGATACACATTCACTGAATACAACATGCACTGGGTGAAACAGAGCCATGGAAAGAGCCTTGAGTGGATTGGAGGTATTA}$ ▶ G Y T F T E Y N M H W V K Q S H G K S L E W I G G I ATCCTAACAATGGTGGTACTAGTTACAACCAGAAGTTCAAGGCCAAGGCCACATTGACTGTAGACAAGTCCTCCAGCAC ▶N P N N G G T S Y N Q K F K A K A T L T V D K S S S T AGCCTACATGGAGCTCCGCAACCTGACATCTGAGGATTCTGCAGTCTATTACTGTGCAAGGGGGGTTTATGATGGTTA A Y M E L R N L T S E D S A V Y Y C A R G V Y D G Y SLLTTGAKAPLSQSPQPKQQPHRSIH TGGCCCCTG ▶ W P L

LIGHT AND FLUORESCENT MICROSCOPE IMAGES CHIMERIC ANTIBODY TEDW BINDING TO S. MUTANS





S. mutans +TEDW

Light microscope

S. mutans +TEDW +Sigma F9512

S. mutans +TEDW +Sigma F5387

Fluorescent microscope